

## Supplemental Figure and Tables

**Table S1. Influence of mucin on MIC.** Liquid MIC results done in SCFM with 0.4% mucin and without mucin grown at 37°C overnight with an inoculum size of 2-7 x 10<sup>5</sup>cells (n=3-5).

| Antibiotic      | MIC ( $\mu\text{g/ml}$ ) |         |
|-----------------|--------------------------|---------|
|                 | + mucin                  | - mucin |
| Gentamicin      | 4                        | 1       |
| Tobramycin      | 2                        | 2       |
| Amikacin        | 16                       | 4       |
| Imipenem        | 0.325                    | 0.625   |
| Meropenem       | 0.125                    | 0.125   |
| Ceftazidime     | 31.25                    | 31.25   |
| Aztreonam       | 4                        | 8       |
| Piperacillin    | 4                        | 4       |
| Erythromycin    | 500                      | 250     |
| Clarithromycin  | 2000                     | 2000    |
| Polymyxin B     | 16                       | 16      |
| Colistin        | 16                       | 2       |
| Norfloxacin     | 16                       | 16      |
| Ciprofloxacin   | 1                        | 0.5     |
| Trimethoprim    | 128                      | 128     |
| Tetracycline    | 64                       | 256     |
| Chloramphenicol | 32                       | 16      |

**Table S2. Resistome mutant susceptibility under surfing conditions - raw data.** Average zone of inhibition measurements for resistome mutants tested against five selected antibiotics. Mutants of up-regulated resistome genes were tested against 10 µg/disk of antibiotic and down-regulated against 100 µg/disk. Statistical significance relative to wild-type was determined using two-way ANOVA. (n=3 resistome mutants; n=6 wild-type) \* p<0.5, \*\* p<0.01, \*\*\* p< 10<sup>-3</sup>, \*\*\*\* p<10<sup>-4</sup>. Standard deviations range from 0 to 2.5mm.

| Mutant                                      | Zone of Inhibition (mm) |              |             |            |             |
|---|-------------------------|--------------|-------------|------------|-------------|
|   | Imipenem                | Tetracycline | Polymyxin B | Tobramycin | Norfloxacin |
| <b>10 µg/disk antibiotic concentration</b>  |                         |              |             |            |             |
| Wild-type                                   | 5.7                     | 5.0          | 5.6         | 3.3        | 1.0         |
| <i>ΔrecG</i>                                | 7.3                     | 8.7*         | 9.7**       | 12.5****   | 7.3****     |
| <i>ΔddaH</i>                                | 9.0*                    | 0***         | 5.3         | 3.0        | 2.3         |
| <i>ΔPA5130</i>                              | 10**                    | 0***         | 0****       | 0*         | 6.5****     |
| <i>cycH</i>                                 | 4.5                     | 4.3          | 11****      | 9.7****    | 0           |
| <b>100 µg/disk antibiotic concentration</b> |                         |              |             |            |             |
| Wild-type                                   | 12.3                    | 6.7          | 8           | 12         | 14.7        |
| <i>ΔarmR</i>                                | 0****                   | 0****        | 1****       | 6.3****    | 0****       |
| <i>ΔPA3576</i>                              | 12.0                    | 3.0*         | 6.0         | 8.3*       | 10.7*       |
| <i>ΔPA1428</i>                              | 12.7                    | 7.7          | 8.0         | 7.0***     | 0.0****     |
| <i>ΔPA2047</i>                              | 12.3                    | 7.0          | 5.7         | 7.3**      | 9.7***      |
| <i>ΔPA1553</i>                              | 9.0                     | 7.3          | 7.7         | 8.0*       | 10.5*       |
| <i>ΔatpB</i> *                              | 9.7                     | 4.0          | 8.0         | 8.3*       | 9.7***      |
| <i>ΔPA4292</i>                              | 7.7**                   | 5.3          | 17****      | 5****      | 10**        |
| <i>ΔclpS</i>                                | 8.3*                    | 6.3          | 15****      | 6.7***     | 8.3****     |
| <i>ΔnuoB</i>                                | 10.7                    | 0****        | 18****      | 6.3****    | 10.3**      |
| <i>ΔPA3721</i>                              | 10                      | 2**          | 14.5****    | 0****      | 10**        |
| <i>ΔPA4429</i>                              | 11                      | 8.7          | 20****      | 0****      | 7.7****     |
| <i>ΔetfA</i>                                | 12.3                    | 9            | 15****      | 7.3**      | 10**        |
| <i>ΔnuoG</i>                                | 9.7                     | 6.5          | 6           | 0****      | 7.3****     |
| <i>ΔPA4781</i>                              | 12.3                    | 10           | 10.3        | 6.7***     | 9****       |
| <i>ΔserA</i>                                | 14.7                    | 8.7          | 12.7**      | 7.3**      | 11*         |
| <i>ΔccmF</i>                                | 14                      | 8.7          | 13.3***     | 7.3**      | 8.7****     |
| <i>ΔPA3667</i>                              | 15.7                    | 0.0****      | 7.7         | 10.0       | 12.0        |
| <i>ΔPA1513</i>                              | 10.7                    | 0.0****      | 7.3         | 10.7       | 14.0        |
| <i>ΔpchF</i>                                | 9.7                     | 5.0          | 7.3         | 1.1***     | 13.3        |
| <i>Δrph</i>                                 | 13.0                    | 9.0          | 6.0         | 5.0****    | 15.0        |
| <i>ΔPA2566</i>                              | 11.0                    | 4.0          | 7.0         | 9.3        | 9.7***      |
| <i>ΔgidA</i>                                | 9.7                     | 7.3          | 9.0         | 10.5       | 10.0**      |
| <i>ΔmutS</i>                                | 16.0                    | 9.3          | 4.3*        | 6.3****    | 11.5        |
| <i>ΔthiG</i>                                | 6.3****                 | 6.7          | 7.0         | 8.7        | 10.3**      |
| <i>ΔnuoF</i>                                | 11.0                    | 5.3          | 6.7         | 7.0***     | 14.7        |
| <i>ΔpckA</i>                                | 9.7                     | 4.8          | 7.0         | 6.3****    | 12.0        |
| <i>ΔPA2571</i>                              | 12.7                    | 6.7          | 7.0         | 7.3**      | 11.7        |

|                 |        |     |     |        |       |
|-----------------|--------|-----|-----|--------|-------|
| $\Delta PA4766$ | 13.7   | 6.0 | 6.5 | 7.0*** | 13.7  |
| $\Delta PA1348$ | 16.7** | 5.5 | 8.3 | 10.7   | 10.7* |
| $\Delta braB$   | 11.0   | 6.7 | 5.7 | 10.0   | 10.7* |
| $\Delta htpX$   | 12.7   | 5.7 | 8.3 | 11.0   | 9.5** |
| $\Delta speA$   | 11.3   | 4.5 | 5.3 | 10.3   | 12.0  |
| $\Delta adhA$   | 12.3   | 7.0 | 6.7 | 10.3   | 13.3  |

**Table S3. RT-qPCR results confirmed the dysregulation of resistome genes shown in RNA-Seq.** The relative fold-change of expression of select resistome genes under surfing conditions (SCFM + 0.4% mucin) relative to swimming (SCFM 0.3% agar) from both the RNA-Seq experiment and RT-qPCR of cells collected from the centre and edge of a surfing colony relative to swimming cells. (FC cut-off of RNA-Seq is  $\pm$  1.5).

|               | Gene Expression (FC) |        |        |      |
|---------------|----------------------|--------|--------|------|
|               | RT-qPCR              | RNASeq | Centre | Edge |
| Gene          | Centre               | Edge   | Centre | Edge |
| <i>recG</i>   | 16.4                 | 8.1    | 1.9    | 2.1  |
| <i>PA5130</i> | 2.2                  | 1.5    | NC     | 2.4  |
| <i>ddaH</i>   | 2.8                  | 3.2    | 4.9    | 2.4  |
| <i>PA1428</i> | -2.2                 | -3.3   | -3.4   | NC   |
| <i>PA2047</i> | 3.1                  | -1.4   | NC     | -2.1 |
| <i>thiG</i>   | -2.5                 | 1.8    | -2.9   | NC   |
| <i>PA3667</i> | -1.2                 | -3.4   | -1.7   | -2.5 |
| <i>PA3576</i> | 2.3                  | -4.4   | NC     | -2.9 |
| <i>atpB</i>   | -1.5                 | 1.1    | -2.1   | NC   |
| <i>PA4292</i> | -2.3                 | -2.4   | -6.7   | NC   |
| <i>muoB</i>   | -1.0                 | -1.2   | -2.8   | NC   |
| <i>PA3721</i> | -2.0                 | 2.1    | -5.3   | -2.7 |
| <i>clpS</i>   | 5.0                  | -3.8   | NC     | -2.3 |
| <i>armR</i>   | -4.4                 | 2.9    | -3.2   | -5.1 |
| <i>cycH</i>   | 5.7                  | 1.4    | NC     | 2.2  |

**Table S4. Primers used in this study.** Primers used for PCR amplification are denoted as F (forward) and R (reverse). All primers used for quantitative PCR (qPCR) are denoted as qF (forward) and qR (reverse).

| Primer Name | Sequence (5'-3')        |
|-------------|-------------------------|
| armR_F      | AGCGTGGCGCCGGAC         |
| armR_R      | CTGCGCGATTCTGATAGCTCA   |
| armR_qF     | CTGAACACTCCCGCAACAA     |
| armR_qR     | GTGTAATCCCCGCCGACCGAC   |
| atpB_F      | GCTTCGGGTTACATCCAGCA    |
| atpB_R      | CGGCGATAGCAGTCAGTCC     |
| atpB_qF     | TGAAGATGAAGGCCTGCAGG    |
| atpB_qR     | TCGGTAGCGGAATGTTCTG     |
| clpS_F      | TTGCAGTTAGTCAAGGCGCT    |
| clpS_R      | AAGGCGAGATTGAGGGTGAC    |
| clpS_qF     | TCAATCAGGACCATCCCGAG    |
| clpS_qR     | AACGGTGGCGGCTGTAATAC    |
| cycH_qF     | TAGACAACGGTTGCCCGT      |
| cycH_qR     | CAGCTAAGCTCAGCGGATT     |
| ddaH_F      | CGATTATCAAGGATTCCAAGGCA |
| ddaH_R      | CCTCCGAAACTGCATCCG      |
| ddaH_qF     | ATCATCGAGGAGACCGTGCA    |
| ddaH_qR     | TGGTCGCCGACCATCATGAT    |
| nuoB_qF     | CGCGTTTCAGGTCTTCTGG     |
| nuoB_qR     | TGCTGCTGCAGGAATCCATC    |
| PA1428_F    | GCGACTGAAGCGTACATGCC    |
| PA1428_R    | ACTTCACCCGTTCTTCAAGC    |
| PA1428_qF   | TTCTGCTTCTGCAACTCGGC    |
| PA1428_qR   | AAGATCCGCCGCTACATTCC    |
| PA2047_F    | CTTGCCCATTCTGTGCCGAT    |
| PA2047_R    | TGTATTGCAAGCGCCCCC      |
| PA2047_qF   | AAGGTGCCGATTCCACGTC     |
| PA2047_qR   | AACGCTACCTGGTATGTCG     |
| PA3576_F    | CCGTCCCGACTTGTCCCG      |
| PA3576_R    | CGAAACGCCACCTTGTGC      |
| PA3576_qF   | TTGCAGGTTGCCGACCAGTT    |
| PA3576_qR   | ATCGGCTATGCCGTGCTGAA    |
| PA3667_F    | ACCCTTCCCCGTAAACCGT     |
| PA3667_R    | TGTTCCCAACTGGCGGTG      |
| PA3667_qF   | GAAGCAGGTTGGCATGGTGT    |
| PA3667_qR   | ACAGAGGCGATCAACCTGGT    |
| PA3721_F    | GCTCTCAGCAAACCGCCT      |
| PA3721_R    | CGAGATCCACCTCACCGATCT   |

|           |                       |
|-----------|-----------------------|
| PA3721_qF | TCCAGGTAGCAGGCGATGAT  |
| PA3721_qR | AAGGCCGACCTGTTCCCTCAA |
| PA4292_qF | GTTCGATCTTCAGCGGGC    |
| PA4292_qR | CGTGGAAAGCCGTTGATGAAC |
| PA5130_qF | ATGGCGTCGACCAGGATGAT  |
| PA5130_qR | ACATCGTCGACGCGCTGAAC  |
| recG_F    | GTCCACCGCGCCATCGAA    |
| recG_R    | CAAGACTGAGACCTACGCCG  |
| recG_qF   | CTGGAGATCTACCATCCGGA  |
| recG_qR   | TGGGTGAGTCCTCGGTAGT   |
| thiG_F    | GATCGTCCACGCCATCGG    |
| thiG_R    | CGCCCCAGGTAGTCGGTAT   |
| thiG_qF   | TGAGCCAAGCATCCAGTACC  |
| thiG_qR   | GTCTCGTCGAGGTCCTTGTA  |

**Table S5. Concentrations of the antibiotics in the disk diffusion assay including their solvents.** Ten  $\mu$ L of antibiotics was added per disk and dried prior to application onto agar surfaces. MeOH – methanol. DMSO – dimethyl sulfoxide.

| Antibiotic      | Concentration ( $\mu$ g/disk) | Solvent |
|-----------------|-------------------------------|---------|
| Gentamicin      | 10                            | Water   |
| Tobramycin      | 10                            | Water   |
| Amikacin        | 5                             | Water   |
| Imipenem        | 10                            | Water   |
| Meropenem       | 5                             | Water   |
| Ceftazidime     | 5                             | Water   |
| Aztreonam       | 30                            | Water   |
| Erythromycin    | 1000                          | MeOH    |
| Clarithromycin  | 500                           | Water   |
| Polymyxin B     | 10                            | Water   |
| Colistin        | 10                            | Water   |
| Norfloxacin     | 5                             | Water   |
| Ciprofloxacin   | 10                            | Water   |
| Trimethoprim    | 1000                          | DMSO    |
| Tetracycline    | 10                            | MeOH    |
| Chloramphenicol | 5                             | Water   |

**Fig S1. Disk diffusion assay plate set-up.** Mid-log phase ( $OD_{600}=0.4-0.5$ ) cultures are inoculated at 1uL around the antibiotic disk at equal distances. Four-point inoculation was used for swim and surf antibiotic disk assays. Disk diffusion control assays were done using a bacterial lawn spread with 50uL of mid-log phase culture and dried antibiotic disks were applied to the centre of each plate.

